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1: *Nucleic Acids Res* 1997 Jan 1;25(1):217-21

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The PROSITE database, its status in 1997.

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The PROSITE database consists of biologically significant patterns and
profiles formulated in such a way that with appropriate computational tools
it can help to determine to which known family of protein (if any) a new
sequence belongs, or which known domain(s) it contains.

MeSH Terms:
- Amino Acid Sequence*
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Substances:
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